

## XPO1 MUTATIONS DRIVE BCR HYPERACTIVATION AND EARLY DISEASE PROGRESSION IN CHRONIC LYMPHOCYTIC LEUKEMIA

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**Introduction:** *XPO1* encodes exportin-1, a key shuttle protein regulating nuclear-cytoplasmic transport. The recurrent hotspot mutation E571K has been linked to early disease progression in chronic lymphocytic leukemia (CLL). ATAC- and RNA-seq analyses revealed that *XPO1*-mutated CLL displays a distinct chromatin landscape with increased accessibility at transcription factor binding sites downstream B-cell receptor (BCR) and upregulation of *MIR155HG*. We hypothesize that mutant *XPO1* enhances nuclear export of *DUSP1*, a nuclear phosphatase inhibiting MAPKs, leading to *MIR155HG* overexpression, increased miR-155 and reduced *SHIP1*, with consequent BCR amplification. This mechanism may be linked to the aggressive clinical behavior and shorter time to first treatment (TTFT) observed in *XPO1*-mutated CLL, although the precise molecular mechanisms have not been fully understood.

**Methods:** To dissect the downstream effects of the E571K, HG3 CLL cell line was engineered using CRISPR-Cas9. A control was generated in parallel. Clones were validated by next generation sequencing and analyzed by flow cytometry for immunophenotype, phosphoflow and calcium flux upon anti-IgM/IgD stimulation. *XPO1*-target protein colocalization was assessed by ImageStream and Western blot.

**Results:** HG3 CLL cell line was engineered through CRISPR-Cas9 to introduce E571K mutation together with two additional silent mutations disrupting the Cas9 recognition site, pre-

venting further cleavage after repair. In parallel, a control cell line harboring only the two silent mutations was generated. All CRISPR-edited clones retained a CLL-like immunophenotype with lambda light chain restriction and surface IgM/IgD positivity. Both the *XPO1*<sup>E571K</sup> and control clones carried homozygous del13q and wild-type *TP53*. To evaluate BCR reactivity, calcium flux was measured. The E571K clone displayed stronger calcium flux upon anti-IgM stimulation ( $p=0.009$ ) compared to the control (Figure 1A) and a similar trend was observed upon anti-IgD stimulation. This is possibly explained by higher phosphorylated levels of p-SYK ( $p<0.0001$ ), p-BTK ( $p<0.0001$ ), p-AKT ( $p=0.009$ ), p-ERK ( $p=0.0001$ ) compared to the control, suggesting a more active BCR signaling (Figure 1B-E). Furthermore, *XPO1*<sup>E571K</sup> cells showed increased cytoplasmic colocalization of *DUSP1* with *XPO1*, consistent with enhanced nuclear export. In addition, under serum withdrawal, *XPO1*-mutated cells showed reduced cell death at 72 hours compared to control, indicating increased survival under stress conditions.

**Conclusions:** Preliminary *in vitro* investigations utilizing *XPO1*-mutant engineered HG3 CLL cell lines demonstrate enhanced BCR signaling activity, corroborating prior transcriptomic (RNA-seq) and chromatin accessibility (ATAC-seq) analyses. Ongoing studies are focused on delineating the specific *XPO1* cargo proteins that mediate this enhanced BCR signaling phenotype.

CHRONIC LYMPHOCYTIC LEUKEMIA AND CHRONIC LYMPHOPROLIFERATIVE DISORDERS

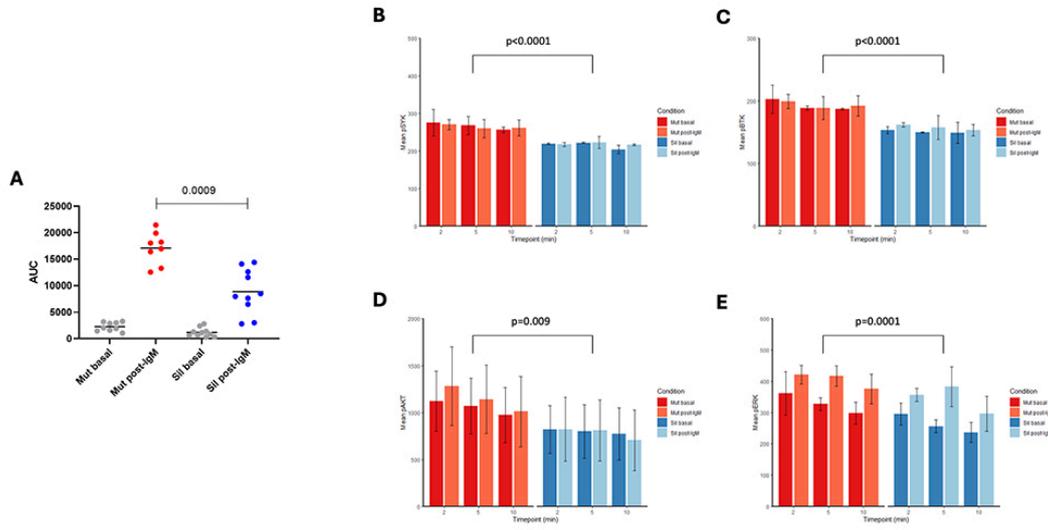


Figure 1